Attorney's Docket No.: 14875-137US1 / C1-A0210Y1P-US

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REMARKS

The amendments to the specification clarify the translated text, correct errors, and, identify abbreviations and trademark information in the original Japanese specification, all of which are apparent when read in the context of the entire specification. The amendments also correct the title and insert into the specification a paper copy of the sequence listing in which the general information has been updated to reflect accurate information for the instant application. In addition Fig. 1 is amended to move the line denoting the EcoR1 restriction enzyme recognition site from after the "t" to after a "c."

Claims 1-20 are pending. Claim 10 is amended to remove multiple dependency. New claim 20 is based on original claim 10. Applicants ask that all claims be examined in view of the amendment to the claims. No new matter has been introduced by these amendments.

Please apply any other charges to deposit account 06-1050, referencing Attorney-Docket No. 14875-137US1.

Respectfully submitted,

Date: December 3, 2004

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Amendments to the Drawings:

The attached replacement sheet of drawings includes changes to Fig. 1 and replaces the original sheet including Fig. 1.

In Figure 1, the line denoting the EcoR1 restriction enzyme recognition site has been moved from after the "t" to after a "c."

Attachments following last page of this Amendment:

Replacement Sheet (1 page)
Annotated Sheet Showing Change (1 page)

Page 1 of 9 Appl. No.: 14875-137US1 Annotated Sheet Showing Change(s)

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												g	aat	toc Ser	acc'	0
atg Met	SEO gta Val SEO	agc Ser	gct Ala	att Ile	gtt Val	tta Leu	tat Tyr	gtg Val	ctt Leu	t t g Leu	gcg Ala	gcg Ala	gcg	gcg	cat	 48
64F1/ SEQ ID NO: 1																
tct	gcc	ttt	gcg	gcg	gag	cac	tgc	aac	gcg	caa	atg	aag	acg	ggt	ccg	96
Ser	Ala	Phe	Ala	Ala	Glu	His	Cys	Asn	Ala	Gln	Met	Lys	Thr	Gly	Pro	
tac	aag	att	aaa	aac	ttg	gac	att	acc	ccg	ccc	aag	gaa	acg	ctg	caa	144
Tyr	Lys	Ile	Lys	Asn	Leu	Asp	Ile	Thr	Pro	Pro	Lys	Glu	Thr	Leu	Gln	
aag	gac	gtg	gaa	atc	acc	atc	gtg	gag	acg	gac	tac	aac	gaa	aac	gtg	192
Lys	Asp	Val	Glu	Ile	Thr	Ile	Val	Glu	Thr	Asp	Tyr	Asn	Glu	Asn	Val	
att	atc	ggc	tac	aag	ggg	tac	tac	cag	gcg	tat	gcg	tac	aac	ggc	ggc	240
Ile	Ile	Gly	Tyr	Lys	Gly	Tyr	Tyr	Gln	Ala	Tyr	Ala	Tyr	Asn	Gly	Gly	
tcg	ctg	gat	ccc	aac	aca	cgc	gtc	gaa	gaa	acc	atg	aaa	acg	ctg	aat	288
Ser	Leu	Asp	Pro	Asn	Thr	Arg	Val	Glu	Glu	Thr	Met	Lys	Thr	Leu	Asn	
gtg	ggc	aaa	gag	gat	ttg	ctt	atg	tgg	agc	atc	agg	cag	cag	tgc	gag	336
Val	Gly	Lys	Glu	Asp	Leu	Leu	Met	Trp	Ser	Ile	Arg	Gln	Gln	Cys	Glu	
gtg	ggc	gaa	gag	ctg	atc	gac	cgt	tgg	ggc	agt	gac	agc	gac	gac	tgt	384
Val	Gly	Glu	Glu	Leu	Ile	Asp	Arg	Trp	Gly	Ser	Asp	Ser	Asp	Asp	Cys	
t t t	cgc	gac	aac	gag	ggc	cgc	ggc	cag	tgg	gtc	aaa	ggc	aaa	gag	ttg	432
Phe	Arg	Asp	Asn	Glu	Gly	Arg	Gly	Gln	Trp	Val	Lys	Gly	Lys	Glu	Leu	
gt g	aag	cgg	cag	aat	aac	aat	cac	ttt	gcg	cac	cac	acg	tgc	aac	aaa	480
Val	Lys	Arg	Gln	Asn	Asn	Asn	His	Phe	Ala	His	His	Thr	Cys	Asn	Lys	
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Ser	Trp	Arg	Cys	Gly	Ile	Ser	Thr	Ser	Lys	Met	Tyr	Ser	Arg	Leu	Glu	
t gc	cag	gac	gac	acg	gac	gag	tgc	cag	gta	tac	att	ttg	gac	gct	gag	576
Cys	Gln	Asp	Asp	Thr	Asp	Glu	Cys	Gln	Val	Tyr	Ile	Leu	Asp	Ala	Glu	
ggc	aac Asn	ccc Pro	atc Ile	aac Asn	gtg Val	acc Thr	gtg Val	gac Asp	ac t Thr	gtg Val	ctt Leu	cat His	cga Arg	gac Asp	ggc Gly	624
gtg	agt	atg	at t	ctc	aaa	caa	aag	tct	acg	ttc	acc	acg	cgc	caa	ata	672
Val	Ser	Met	I l e	Leu	Lys	Gln	Lys	Ser	Thr	Phe	Thr	Thr	Arg	Gln	Ile	
aaa	gct	gcg	tgt	ctg	ctc	att	aaa	gat	gac	aaa	aat	aac	ccc	gag	tcg	720
Lys	Ala	Ala	Cys	Leu	Leu	Ile	Lys	Asp	Asp	Lys	Asn	Asn	Pro	Glu	Ser	
gtg	aca	cgc	gaa	cac	tgt	t t g	att	gac	aa t	gat	ata	tat	gat	ctt	tct	768
Val	Thr	Arg	Glu	His	Cys	Leu	Ile	Asp	Asn	Asp	Ile	Tyr	Asp	Leu	Ser	
aaa	aac	acg	tgg	aac	t gc	aag	ttt	aac	aga	tgc	att	aaa	cgc	aaa	gtc	816
Lys	Asn	Thr	Trp	Asn	Cys	Lys	Phe	Asn	Arg	Cys	Ile	Lys	Arg	Lys	Val	